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                          GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on:         May   4, 2005, 12:02:42 ; Search time 304.335
Seconds
                                           (without alignments)
                                           1814.018 Million cell
updates/sec

Title:          US-09-300-482-311
Perfect score:  274
Sequence:       1
cttacattcctttctccacc..........gccaagcatgccgtcgaccg 274

Scoring table:  IDENTITY_NUC
                Gapop 10.0 , Gapext 1.0

Searched:       9369276 seqs, 1007425964 residues

Total number of hits satisfying chosen parameters:     18738552

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :       Pending_Patents_NA_New:*
                 1:  /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*
                 2:  /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq2:*
                 3:  /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
                 4:  /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
                 5:  /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
                 6:  /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
                 7:  /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
                 8:  /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq2:*
                 9:  /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq3:*
                 10: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq:*
                 11: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq2:*
                 12: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq3:*
                 13: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:*
```